

FIGURE 1

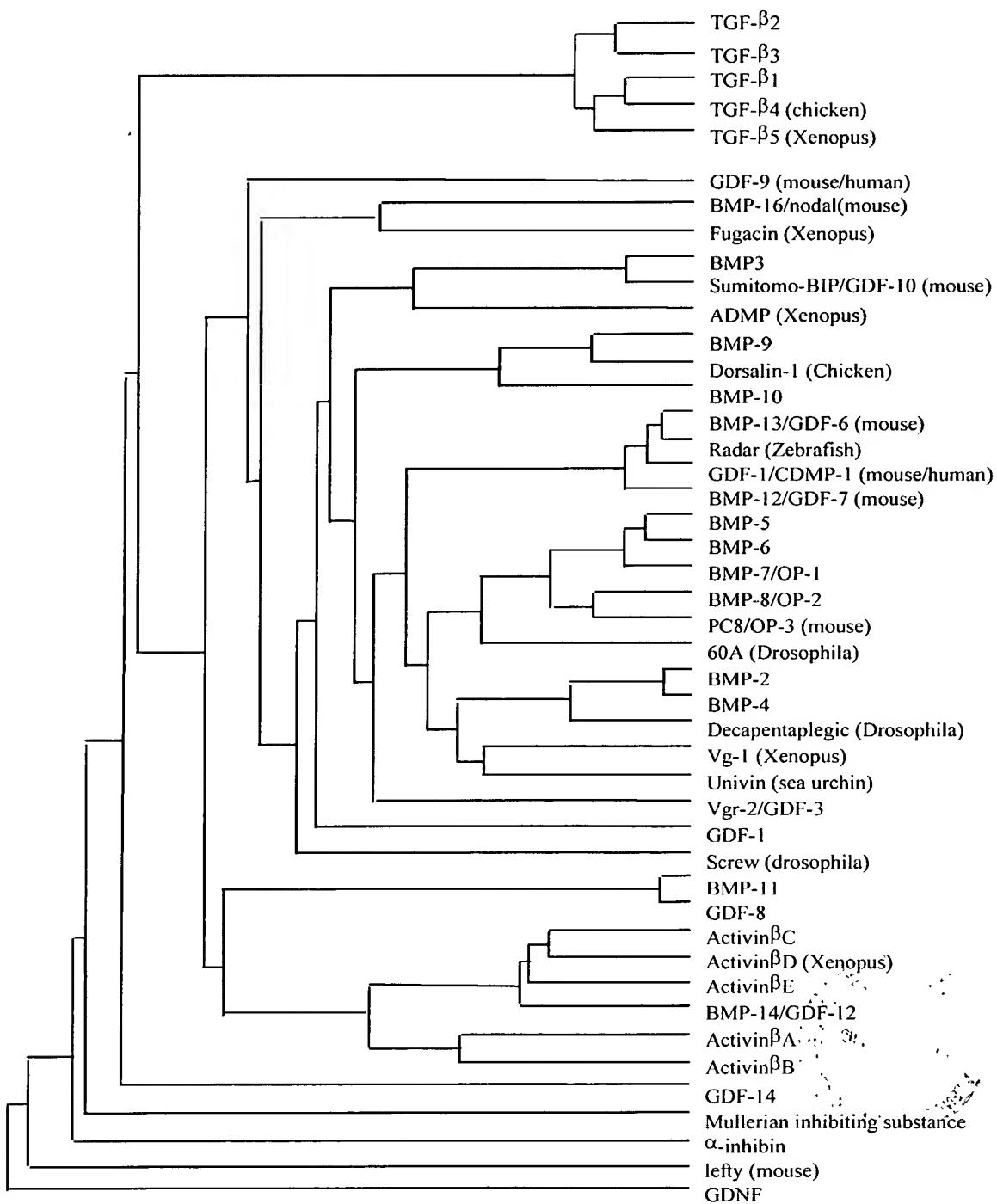


FIGURE 2A

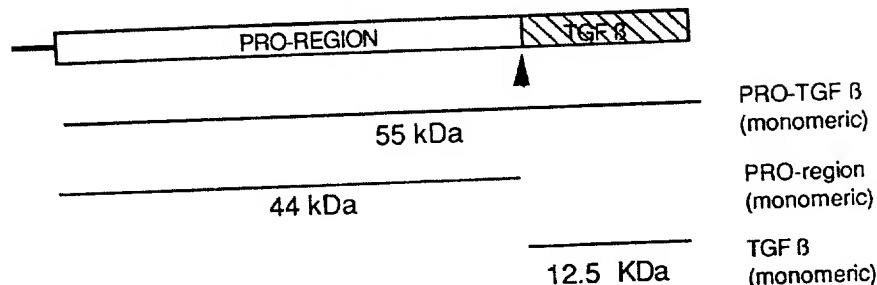


FIGURE 2B

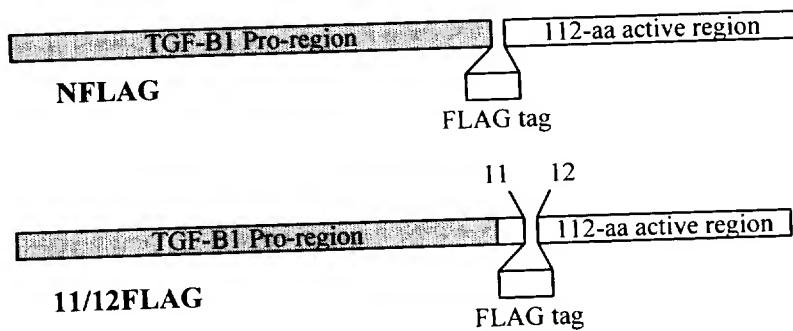


FIGURE 3A

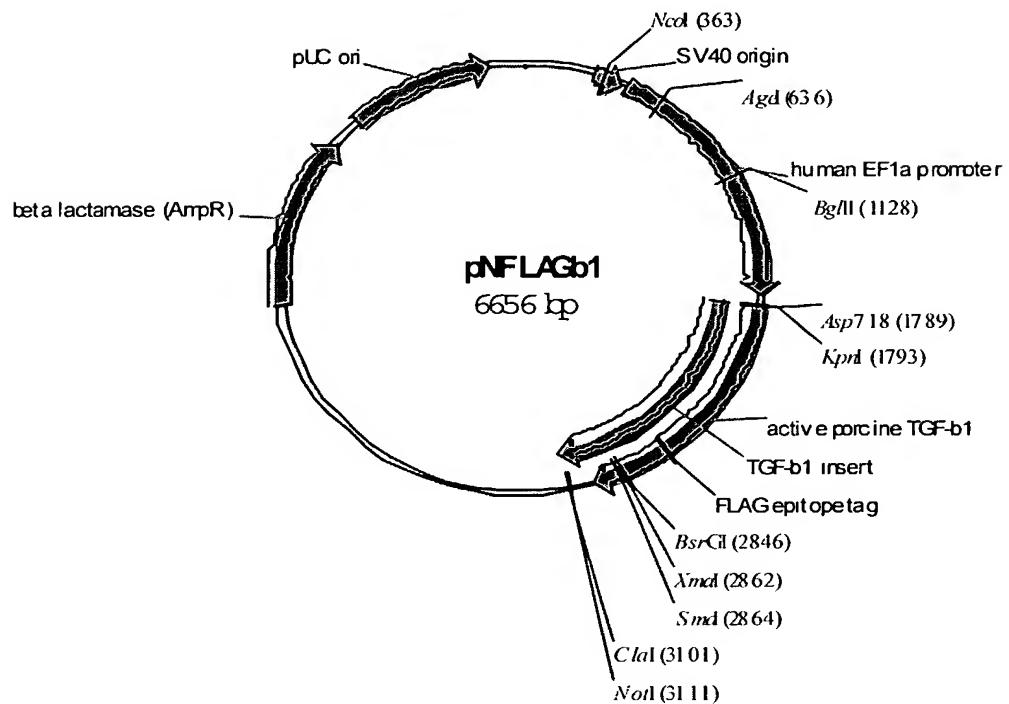


FIGURE 3B

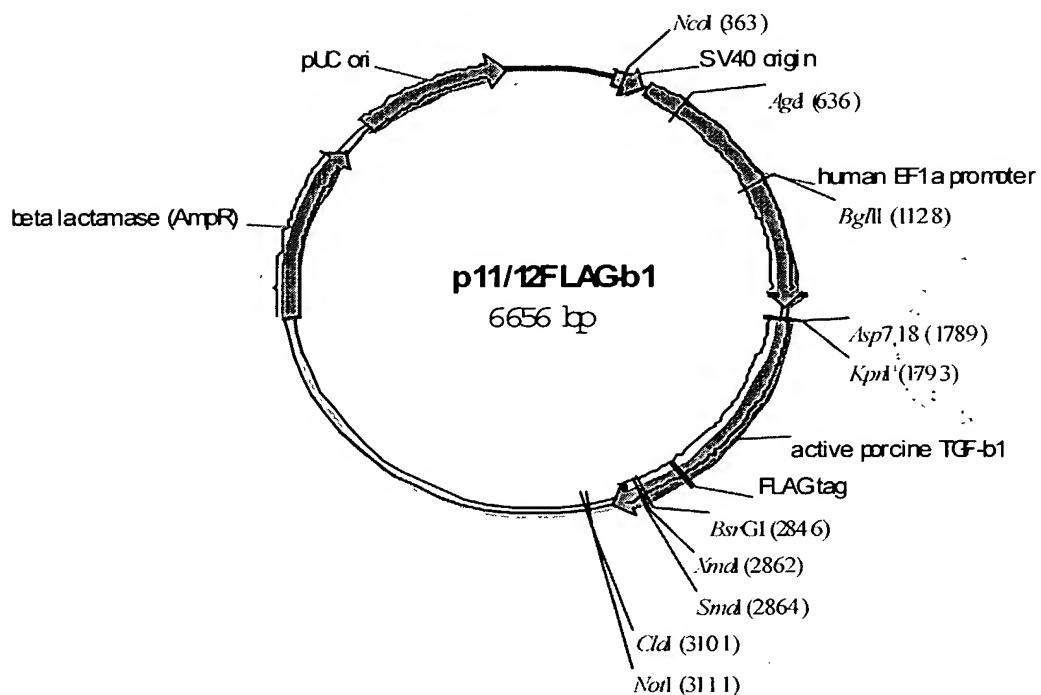


FIGURE 4A

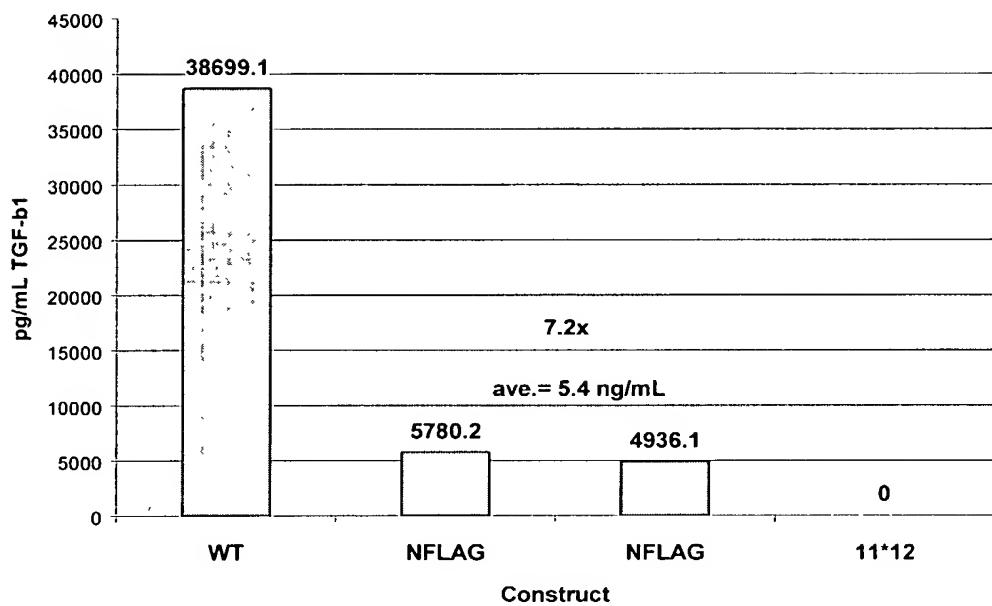


FIGURE 4B



FIGURE 5

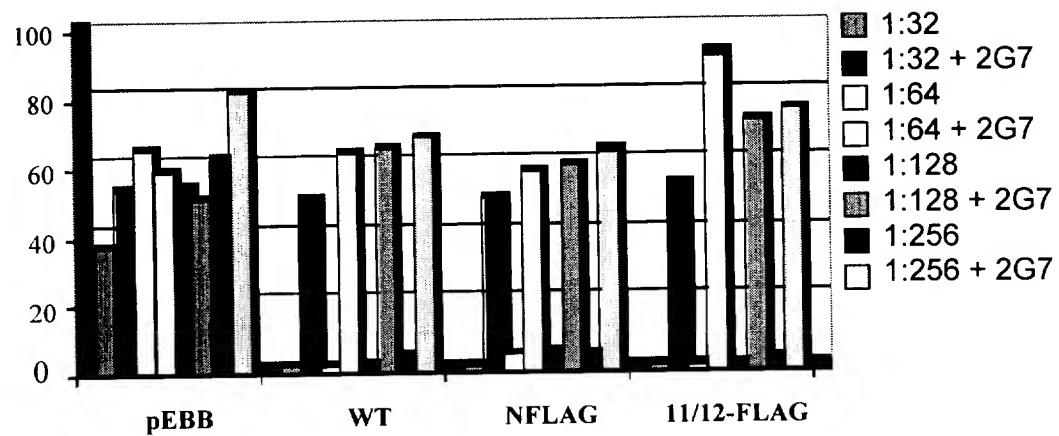


FIG 6A

N-FLAG-TGF- β 1:

2601 S S R H R R ↓ D Y K D D D D
 AGCTCCCG GCACCGCCGA GACTACAAGG ATGACGACGAA
 TCGAGGGC CGTGGCGGCT CTGATGTTCC TACTGCTGCT
 2651 K A L D T N Y C F S S T E K N C
 CAAGGCCCTG GATAACACT ACTGCTTCAG CTCCACGGAG AAAGAACTGCT
 GTTCCGGGAC CTATGGTTGA TGACCGAAGTC GAGGTGCCCTC TTCTTGACGA
 C V R Q L Y I D F R K D L G W K W
 2701 GCGTGGGCA GCTCTACATT GACTTCCGGA AGGACCTGGG CTGGAAGTGG
 CGCACCCGT CGAGATGTA CTGAAGGCCCT CCCTGGACCC GACCTTCACC

FIG 6B

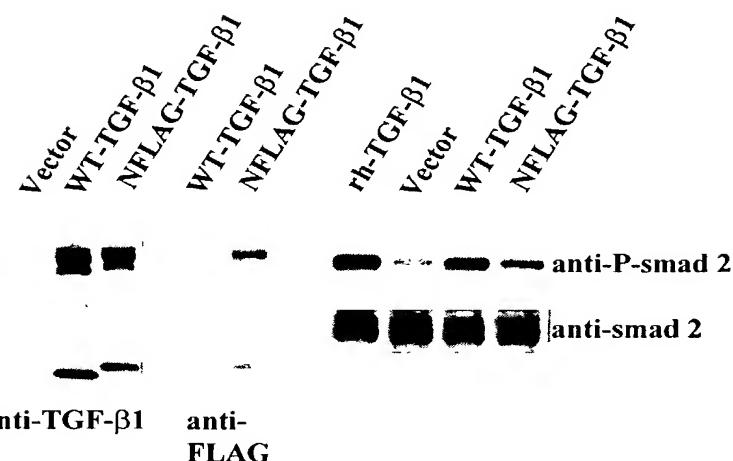


FIG 6C

2601 Q H L H S S R H R R ↓ A L D T N S Y
 CAGCACCTGC ACAGCTCCCG GCACCGCCGA GCGCTGGATA CCAACGACTA
 GTCGTGGACG TGTGGAGGGC CGTGGCGGCT CGGGACCTAT GGTTGTCGAT
 2651 K D D D D K A L D T N Y C F S S
 CAAGGATGAC GACGACAAGG CCTGGATAC CAACTACTGC TTCAAGCTCCA
 GTTCCCTACTG CTGCTGTTC GGGACCTATG GTTGATGAGG AAGTCGAGGT

2601 Q H L H S S R H R R ↓ A L D T N S Y
 CAGCACCTGC ACAGCTCCCG GCACCGCCGA GCGCTGGATA CCAACGACTA
 GTCGTGGACG TGTGGAGGGC CGTGGCGGCT CGGGACCTAT GGTTGTCGAT
 2651 P Y D V P D Y A S L A L D T N Y
 CCCATACGAC GTGCCAGACT ACGCATCTCT GCGCTGGAT ACCAACTACT
 GGGTATGCTG CACGGTCTGA TGGTAGAGA CGGGACCTA TGGTTGATGA

FIG 6D

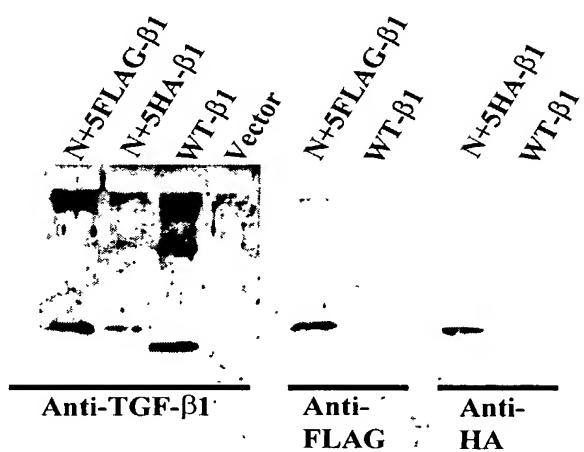


FIG 7A

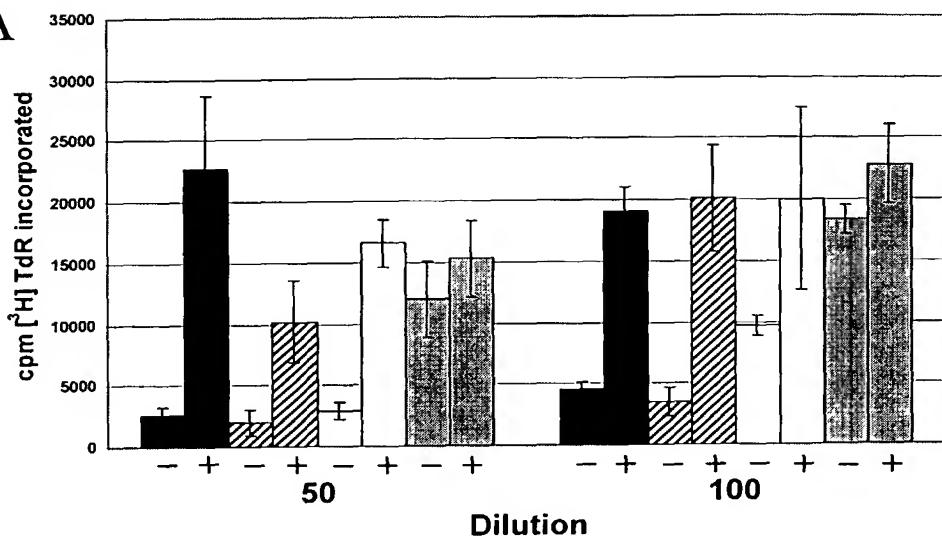


FIG 7B

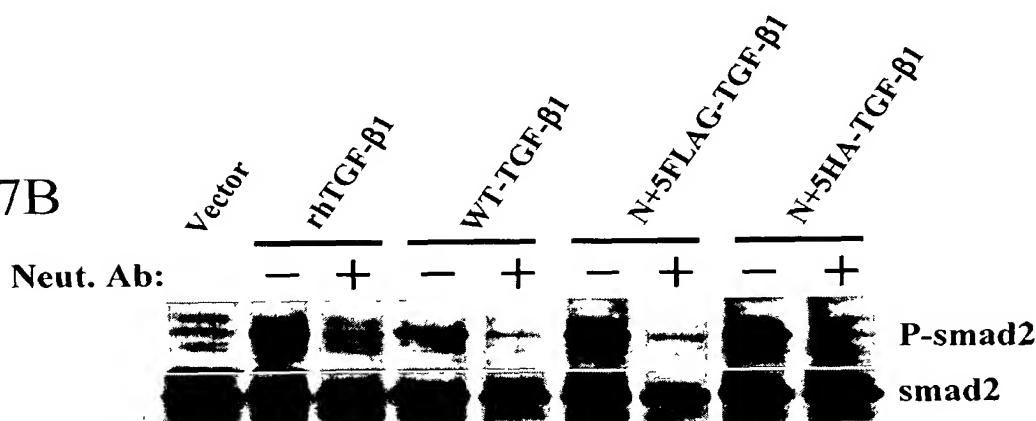


FIG 7C

